



SEQUENCE LISTING

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<120> LOOP PROBE HYBRIDIZATION ASSAY FOR POLYNUCLEOTIDE ANALYSIS

<130> 5100-0703.01

<140> 10/716,063

<141> 2003-11-17

<150> 09/815,510

<151> 2001-03-22

<150> 60/191,227

<151> 2000-03-22

<150> 60/237,000

<151> 2000-09-29

<160> 24

<170> PatentIn version 3.2

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<222> (23)..(23)

<223> N = (biotin dT)

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<223> N = (biotin dT)

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27

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27

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<400> 10
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 cccagtgtgg aagaggacca ttcctctgg g 31

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 <400> 13
 tcacaggttc cgatgtcaac 20

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 <400> 14
 cagggtggtc ctctcacac 19

 <210> 15
 <211> 151
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 <220>
 <223> Sequence spanning the D7S8 locus upper strand

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 <221> misc_feature
 <222> (50)..(50)
 <223> N: Location of SNPN=C/G: Allele AN=T/A: Allele B

<400> 15
 ctagggatgt tcctgtctca gggaccctga ccttattgct cccctttccn ggaatgctgg 60
 tcctgacaca ataataaag ctctgagaag gcagccattt ttgtatgctt tactccaggc 120
 tactttctcaa ctcgcagaac agggcttggc a 151

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 <223> Sequence spanning the D7S8 locus lower strand

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 <221> misc_feature
 <222> (102)..(102)
 <223> N: location of SNPN=C/G: Allele AN=T/A: Allele B

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 ccttctcaga gcttatatta ttgtgtcagg accagcattc cnggaaaggg gagcaataag 120
 gtcagggtcc ctgagacagg aacatcccta g 151

<210> 17
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 <212> DNA
 <213> Artificial

<220>
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<400> 17
 gggaccctga ccttattgc 19

<210> 18
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 <212> DNA
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<220>
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tggtgcctt ctcagagct 19

<210> 19
<211> 21
<212> DNA
<213> Artificial

<220>
<223> 151bp fragment PCR forward primer

<400> 19
ctagggatgt tcctgtctca g 21

<210> 20
<211> 20
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<223> 151bp fragment PCR reverse primer

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tgccaagccc tggtctgcga 20

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<220>
<223> mbD7S8B

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ttttttacca gcattccggg aaag

24

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<400> 24
ttttttacca gcattccagg aaag

24